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High viral loads despite absence of clinical and pathological findings in cats experimentally infected with feline coronavirus (FCoV) type I and in naturally FCoV-infected cats

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Accepted 15 August 2003

Summary Specified pathogen-free cats were naturally infected with FCoV or experimentally infected with FCoV type I. Seroconversion was determined and the course of infection was monitored by measuring the FCoV loads in faeces, whole blood, plasma and/or monocytes. Tissue samples collected at necropsy were examined for viral load and histopathological changes. Experimentally infected animals started shedding virus as soon as 2 days after infection. They generally displayed the highest viral loads in colon, ileum and mesenteric lymph nodes. Seroconversion occurred 3–4 weeks post infection.

Naturally infected cats were positive for FCoV antibodies and monocyte-associated FCoV viraemia prior to death. At necropsy, most animals tested positive for viral shedding and FCoV RNA was found in spleen, mesenteric lymph nodes and bone marrow. Both experimentally and naturally infected cats remained clinically healthy. Pathological findings were restricted to generalized lymphatic hyperplasia. These findings demonstrate the presence of systemic FCoV infection with high viral loads in the absence of clinical and pathological signs.

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Introduction

Feline infectious peritonitis (FIP) is a fatal disease of cats, induced by feline coronavirus (FCoV). FIP is currently the leading infectious cause of death in pet cats (Fehr et al., 1997; Pedersen, 1995).

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Until now, the pathogenesis of the disease was not well understood. In 1982 it was postulated that FIP was an immune-mediated disease (Jacobse-Geels et al., 1982). One manifestation of an immune-mediated disease is the antibody-dependent enhancement (ADE) frequently observed in experimental infection (Weiss and Scott, 1981). However, ADE does not appear to be widely occurring under field conditions (Addie et al., 1995; Fehr et al., 1996).

Originally, it was thought that FIP was caused by a virulent feline infectious peritonitis virus (FIPV) (Pedersen, 1976; Ward, 1970). Later, two types of FCoV were identified, one, which is confined to the digestive tract and does not spread beyond regional lymph nodes. This is the so-called feline enteric coronavirus (FECV). The second, immunologically closely related to FECV, is capable of going systemic by infecting monocytes and macrophages and causing FIP. This latter is the so-called FIPV (Pedersen et al., 1981b; Stoddart et al., 1988; Stoddart and Scott, 1989).

Recent findings support the hypothesis that FIPVs arise from avirulent FCoVs (FECVs) due to mutations during their replication (Horzinek and Lutz, 2000; Poland et al., 1996; Vennema et al., 1998). Moreover, new techniques such as RT-PCR could demonstrate the presence of FCoV in the blood of healthy cats (Addie et al., 1996; Egberink et al., 1995; Fehr et al., 1996; Foley et al., 1997; Gunn-Moore et al., 1998; Kipar et al., 1999). These observations support the hypothesis that viraemia is not a hallmark of FIP-inducing FCoVs and that a carrier state exists where cats remain healthy despite being systemically infected. Persistently infected, healthy cats are believed to play the most important epidemiological role in FIP, because they represent a constant source of infection by shedding FCoV in their faeces (Foley et al., 1997).

Two FCoV serotypes have been identified on the basis of *in vitro* neutralization tests (Hohdatsu et al., 1991a,b; Pedersen et al., 1984a), both of which can cause FIP. The type I virus group represents the genuine FCoVs; they grow poorly in cell culture. Type II viruses, however, arise from recombinations between feline and canine (CCV) coronavirus (Herrewegh et al., 1998). These proliferate well in cell culture and are widely used for the study of FCoV infection. However, in the field, serotype II accounts for only 20 to 30% of all FCoV infections at least in Japan (Hohdatsu et al., 1992). In a recent study conducted in Austria, type I was found to be present in 62% of the cats studied (Posch et al., 2001).

The goal of the present study was to quantify viral loads in the intestinal tract and in various organs of cats infected with FCoV type I and to correlate these with potential clinical and pathological signs. As type I viruses are difficult to grow in cell culture, viruses originating from faeces of naturally infected animals were used for experimental studies. As natural infection occurs via the faecal-oral route, animals were inoculated perorally. Furthermore, in a prospective experiment with naturally infected animals, the presence of FCoV in faeces, blood monocytes and different organs was evaluated and correlated with clinical and pathological findings.

Materials and methods

Animals

For experimental infections, 15 specified pathogen-free (spf) kittens were provided by IFFA-Credo (Saint-Germain sur l'Arbresle, France) at either 6 or 16 weeks of age. The animals were first acclimatized by keeping them together for 4 days and were then separated into groups. After infection, the cats were clinically examined daily, weighed, and the rectal temperature was measured.

Eight spf cats were provided by Harlan Winkelmann (Borchen, Germany) at 9 weeks of age. They were acclimatized until they entered the experiment at the age of 21 weeks (Kipar et al., 1999).

Preparation of virus inoculum for infection

Viruses to be used as the source of infection were isolated from faecal samples of field cats or from the intestines of cats experimentally infected with an FCoV field strain. Based on the determination of the sequence of part of the S-gene and on the comparison with sequences deposited in the gene bank, they all belonged to type I.

Faecal samples were prepared as follows: 20 ml of ice-cold RPMI 1640 medium (Sigma, Schnellendorf, Germany) containing L-glutamine and 10% fetal calf serum (FCS, BioConcept, Allschwil, Switzerland), but no antibiotics, were added to 1.5–2 g of faeces under sterile and RNase-free conditions. The samples were incubated for 10 minutes at 4 °C with occasional shaking and finally centrifuged twice for 10 minutes at 900×g to obtain a clear supernatant. The supernatants were aliquoted into tubes of 4 ml each, frozen, and stored at –80 °C. Aliquots of these virus-containing supernatants were analysed

Table 1 Compilation of experimental data of the three infection experiments

Experiment	Group	Cat no.	Virus		Dose (RNA copy no.)
			Strain	Origin	
I	1	3137B	FCoVZu5+FCoVZu2	Faecal extract	$1.3 \times 10^4 + 8.8 \times 10^4$
		3154B			
	2	3132B	FCoVZu1	Faecal extract	0.8×10^6
II	3	3141B	FCoVZu3	Faecal extract	$1.5 \times 10^5 + 1.5 \times 10^5$
		3138B			
	4	3139B	FCoVZu1	Faecal extract	8×10^6
III	5	3312B	FCoVZu1	Gut homogenate	3×10^9
		3313B			
	6	3314B	FCoVZu1	Gut homogenate	0.3×10^9
		3252B			
		3258B			
	7	3257B	FCoVZu1	Gut homogenate	0.03×10^9
		3265B			
		3253B			
		3256B			

to determine the FCoV load. In parallel, the samples were tested for the presence of parvoviruses (see below), yielding negative results. Four of these faecal extracts containing FCoV were used to infect kittens: FCoVZu1, FCoVZu2, FCoVZu3 and FCoVZu5.

The intestines (gut homogenates) were prepared as follows: the intestines including their contents were collected from three kittens that had been experimentally infected with the FCoVZu1 strain. The intestines were collected after euthanasia at peak infection and separated into duodenum, jejunum, ileum and colon. Samples were snap-frozen in liquid nitrogen and stored at -80°C . After it became evident that the viral loads were high in intestinal samples of all three cats, the intestinal aliquots were homogenized and pooled in a total volume of 500 ml of stabilizing solution (HONDEV STAB, Intervet UK; diluted 1:4 in RPMI 1640). Homogenization was performed using an Ultra Tourax instrument (Kinematica AG, Lucerne, Switzerland) at 20,000 rpm for 15 minutes on ice at 5–6 intervals. The final volume of about 500 ml of pooled homogenate was aliquoted and stored at -80°C . This gut homogenate was later used in a titration experiment in cats.

Infection experiments

Experimentally infected cats

Animals were infected perorally twice within 24 hours, the infectious dose was applied via an oesophageal tube. This was done in order to keep the volume of the inoculum as small as possible and to increase the chance of infection. Infection was

performed while the animals were anaesthetized to avoid any unwanted reaction from the cats that could compromise any efforts to standardize the procedure. In the first infection experiment, six 6-week-old spf kittens were divided into three groups of two animals each and infected with different FCoV field strains (Table 1). In the second infection experiment, three spf kittens of 6 weeks of age were kept together as one group during the entire experiment (17 days). Each cat was inoculated perorally with two doses of the FCoVZu1 strain within 24 hours (Table 1). In the third infection experiment, six 16-week-old spf kittens were divided into three groups of two animals each and infected with ten-fold dilutions (see Table 1) of the virus strain FCoVZu1.

Prior to infection, and at weekly intervals, 1 ml of blood was collected from all animals and haematological parameters as well as FCoV viral loads were determined.

FCoV shedding was determined by measuring FCoV viral load in faecal swabs. In the first two infection experiments, faecal swabs were collected on a daily basis, whereas in the third infection experiment shedding was determined during the first 5 weeks in faecal swabs collected every 2 days; in the remaining period, samples were collected weekly.

Naturally infected cats

Eight spf cats, originating from a natural FCoV contact infection experiment in which a total of 20 cats had been included, were housed together with 11 FCoV-infected, clinically healthy cats (tested

positive for circulating and FCoV-specific immune complexes (Kipar et al., 1999) originating from animal shelters, for 30 weeks from the age of 21 weeks on. The eight cats survived the total observation period, whereas the remaining 12 animals of that group developed FIP (Kipar et al., 1999). The cats were clinically examined on a weekly basis throughout the experiment. Blood for serology and monocyte isolation was collected every 2 months. Viral loads and pathological changes were assessed on samples collected at necropsy.

Preparation of viral RNA from whole blood, plasma and faeces

RNA was extracted from whole EDTA blood using the QIAamp[®] RNA Blood Mini Kit (Qiagen, Basel, Switzerland) according to the manufacturer's protocol. For the extraction of viral RNA from plasma the QIAamp[®] Viral RNA Mini Kit (Qiagen) was used.

To determine the viral load in faeces, faecal swabs were inserted into the rectum and then placed in 1.5 ml RNase-free Eppendorf tubes. RNA was prepared as follows: 200 µl of phosphate buffered saline (PBS 1×) were added to each swab, the tubes were closed and briefly mixed by vortexing. The swabs were then incubated at 42 °C for 10 minutes and mixed again. After a short spin the swab was inverted in the tube and another short spin was applied. The dried swab was removed and the extract was inactivated by incubating at 95 °C for 5 minutes. Any residual faecal material was eliminated in a final centrifugation step. The faecal extracts were diluted 1:10 in PBS 1× and both the undiluted and the diluted solution were tested for the presence of FCoV RNA.

Viral RNA isolation from monocytes

As previously described (Kipar et al., 2001b), monocytes were isolated from heparin blood (4.5 ml) using Ficoll (Histopaque[®]-1077, Sigma). The lymphocyte-enriched fraction was resuspended in 3 ml of RPMI 1640 medium supplemented with 10% FCS (BioConcept), antibiotics (Penicillin, Streptomycin) and L-glutamine and the cells were distributed in three wells of a 24-well plate. Cells were incubated for 16 hours in 5% CO₂ to allow monocytes to adhere to the well surface. Floating lymphocytes were washed out. The monocytes of all three wells were lysed together in 350 µl RLT Buffer (Qiagen), the lysate was homogenized over a QIAshredder[®] column (Qiagen) and RNA was subsequently purified using the RNeasy[®] Mini Kit (Qiagen) according to the manufacturer's protocol.

In naturally infected cats isolated monocytes, obtained 4 months after the beginning of the experiment and 2 months later (end of experiment), were co-cultivated with whole feline embryo cells (WFE) and RNA was isolated as described elsewhere (Gunn-Moore et al., 1998).

RNA isolation from organs

In experimentally infected animals, samples from different organs (duodenum, jejunum, ileum, colon, mesenteric lymph nodes, liver, spleen, bone marrow, kidney, thymus, lung, tonsil and brain) were collected under sterile, RNase-free conditions and directly frozen in liquid nitrogen. The tissues were then stored at -80 °C. RNA was purified by means of the ABI Prism[™] 6700 Automated Nucleic Acid Workstation (Applied Biosystems (ABI), Rotkreuz, Switzerland). About 30 mg of tissue was cut out from the frozen samples and placed in a 2 ml Eppendorf tube containing 1.2 ml of ABI Lysis Buffer and a 3 mm steel bead (Schieritz & Hauenstein AG, Arlesheim, Switzerland). The samples were homogenized and lysed by mixing twice for 1 minute at 30 Hz in a Mixer Mill device (Qiagen). After mixing, the samples were incubated for 30 minutes on ice to reduce the foam that had formed during the homogenization step. After equilibration to room temperature (RT), the lysed samples were digested during 1 hour with 200 µg of proteinase K (Qiagen). The samples were then subjected to another round of homogenization with the Mixer Mill and finally stored at -20 °C for no longer than 2 days. From these lysed samples, RNA was purified using the ABI Prism[™] 6700 chemistry according to the manufacturer's protocol. Negative controls containing only PBS 1× were also extracted to check eventual cross-contamination during this extraction step.

As control for the efficacy of RNA purification, the RNA was reverse transcribed using the High Capacity cDNA Archive Kit (Applied Biosystems) by means of the ABI Prism[™] 6700. The expression of the housekeeping gene glyceraldehyde-3-phosphate-dehydrogenase (GAPDH) was measured by real-time PCR (Leutenegger et al., 1999b).

In naturally infected animals, for RNA extraction, approximately 100 mg of tissue was homogenized with a tissue homogenizer (PCR Tissue Homogenizing Kit, Süd-Laborbedarf GmbH, Gauting, Germany) and lysed in each 700 µl lysis buffer. Total RNA was extracted with the RNeasy mini kit (Qiagen, Hilden, Germany) according to the manufacturer's protocol (Kipar et al., 2001b). RNA was dissolved in 30 µl RNase-free water.

Viral loads

FCoV viral loads were determined by one-tube real-time RT-PCR detecting a 102 bp fragment of the conserved FCoV 7b gene (Gut et al., 1999), using an automated fluorometer (TaqMan, ABI 7700, Applied Biosystems) in all samples from experimental infections and tissue samples from naturally infected cats.

The infectious dose was calculated as the RNA copy number present in the volume of the infectious inoculum given to the cats (Table 1).

Blood samples of experimentally infected animals were also tested by one-tube real-time PCR for presence of feline parvovirus (FPV), feline leukaemia virus (FeLV), feline immunodeficiency virus (FIV) and feline herpesvirus-1 (FHV-1) as described elsewhere (Hofmann-Lehmann et al., 2001; Leutenegger et al., 1999a; Vogtlin et al., 2002). In the case of FPV, a sequence in the high conserved VP-2 gene was amplified with the TaqMan® technology using the following primers and probe: PV3294f: 5'-ACTGCATCATTGATGGTTGCA-3'; PV3400r: 5'-GGTATGGTTGGTTCCATGGA-3'; PV3375p: 5'-FAM-CCCAATGTCTCAGATCTCATAGCTGCTGG-6-TAMRA-3'.

Large intestinal content was collected from naturally infected cats at necropsy and tested by conventional RT-PCR (Herrewegh et al., 1995) and electron microscopy (negative staining technique) for FCoV (Kipar et al., 1999).

Serology

Antibody titres in serum were determined by immunofluorescence analysis (IFA) as previously described (Osterhaus et al., 1977). Circulating immune complexes were measured by competitive ELISA as described elsewhere (Schroo, 1994).

Results

Infection experiment I

To determine the infectious potential of different FCoV field strains, three groups of two cats each were infected with three different FCoV field isolates (Table 1).

Prior to infection, all cats were negative for FCoV in both blood and faeces. The faeces of the cats of group 2, infected with the FCoVZu1 strain, turned positive on day 4 post infection (p.i.) and the cats continued to shed high amounts of virus (on average 1×10^7 FCoV RNA copies per g faeces) until euthanasia. The animals of group 1 and 3 did not

shed any virus during the first 14 days after infection. The cats of group 2 were sacrificed 14 days p.i., whereas those of group 1 and group 3 were re-infected on day 14 under the same conditions as in the first inoculation. To determine whether lack of infection was a hallmark of the FCoV strain used or whether it depended of the infectious dose, group 1 was re-infected with a different FCoV strain (FCoVZu2), whereas group 3 was re-infected with the same isolate as the first time (FCoVZu3) (Table 1). All four cats then started shedding FCoV in their faeces from day 2 after re-infection on. FCoV RNA was detected in whole blood in two cats (3154B and 3139B) on days 14 and 21 p.i., in cat 3132B on day 7 p.i. and in cat 3138B on day 21 p.i. FCoV RNA was found in the plasma fraction of cat 3132B on day 14 p.i. Cat 3138B was the only animal that seroconverted (day 28 p.i.). The results are compiled in Table 2.

Monocytes were isolated from blood collected prior to euthanasia and cultivated overnight. From all cats tested only in one animal of group 2 (3132B) FCoV RNA could be detected in the monocytes.

Clinically, the cats showed a short episode of moderate pyrexia at days 2 and 3 p.i. The leukocyte counts were slightly elevated at the beginning of the experiment (17,000–27,000 per μ l), but returned to normal values after 1 week. No further clinical signs were detected.

Blood from all animals was additionally tested for the presence of other viruses, including FPV, FeLV, FIV and FHV-1, prior to infection and during the entire experiment with negative results. The experiment was concluded at day 28 p.i.

Infection experiment II

To obtain FCoV-infected intestines for use as infection material for future experiments, three 6-week-old spf kittens were inoculated with the FCoVZu1 strain (Table 1).

All three cats were clinically healthy during the entire observation period, in spite of the fact that they all started to shed high amounts of FCoV in their faeces 2–3 days after infection and continued to shed virus each day until they were sacrificed (Table 2) on day 17 p.i.

Prior to infection, all three cats were negative for FCoV in blood as well as in faeces. All three animals were positive for viral RNA in the plasma fraction on day 7, and one cat (3314B) also on day 17 p.i. In addition, cat 3313B tested positive in whole blood on day 7. Two cats (3312B and 3313B) transiently seroconverted on day 14. Results are compiled in Table 2.

Table 2 Summary of the results of the infection experiments I–III

Parameters	Experiment I			Experiment II	Experiment III		
	Group 1	Group 2	Group 3	Group 4	Group 5	Group 6	Group 7
Virus strain	FCoVZu5+2	FCoVZu1	FCoVZu3	FCoVZu1		FCoVZu1	
Start of shedding (days p.i.)	2–6 ^a	3–4	2–3 ^a	2–3	2	2	2
Intensity of shedding	+++	++/+++	+++	+++	+++	+++	+++
Frequency of shedding; no. of positive tests (samples tested)	18 (58)	22 (30)	19 (58)	46 (54)	32 (48)	37 (48)	37 (48)
Rectal temperature >39 °C	21 (58)	6 (30)	3 (58)	Never	2 (36)	10 (36)	4 (36)
Clinical signs	None	None	None	None	None	None	None
Viraemia, cell-associated; no. of positive tests (samples tested)	2 (10)	1 (6)	3 (10)	1 (12)	0 (22)	0 (22)	0 (22)
Viraemia, cell-free; no. of positive tests (samples tested)	0 (4)	1 (2)	1 (4)	4 (12)	0 (22)	0 (22)	0 (22)
Seroconversion (days p.i.)	–/–	–/–	28/–	14/14/–	14/22	14/14	14/22
Monocyte-associated viraemia; no. of positive tests (samples tested)	0 (2)	1 (2)	0 (2)	1 (3)	0 (22)	0 (22)	0 (22)

+++= C_T value from 20 to 30. ++= C_T value from 30 to 40. –=never seroconverted.

^a=after re-infection.

Blood from all animals was additionally tested for the presence of other viruses, namely FPV, FeLV, FIV and FHV-1, prior to infection and during the entire experiment with negative results.

Infection experiment III

To determine the optimal infectious dose as well as possible dose-dependent effects, three groups of two animals each were infected with ten-fold dilutions of FCoVZu1-infected gut homogenate (Table 1) derived from the intestines of FCoVZu1-infected cats (infection experiment II).

Regardless of the infectious dose, all animals started shedding 2 days after infection. Intensity and frequency of shedding were comparable in all three groups.

In none of the samples tested FCoV RNA was detected in blood or isolated monocytes. All cats seroconverted 2 to 3 weeks p.i. Results are compiled in Table 2.

With the exception of several one day episodes, when the cats exhibited elevated rectal temperatures (≥ 39 °C), all animals remained clinically healthy during the entire observation period of 80 days.

Naturally infected cats

In order to compare experimental with natural infection, we analysed data from animals that originated from an experiment in which FCoV infection was established under long-term natural

exposure by housing of spf cats with naturally infected cats for 5 months (Kipar et al., 1999).

All eight spf cats that survived healthy showed seroconversion with titres of 1:100 or higher when tested either 2 months or 4 months after first being housed with FCoV-infected cats. At the end of the experiment, three animals were still seropositive (titre $\geq 1:100$; Table 3). In six cats (cat Nos. 2–5, 7, 8), circulating FCoV-specific immune complexes were detected at least at one time point, whereas at the end of the experiment all cats were negative in the competitive ELISA (Kipar et al., 1999).

Four months after first being housed with FCoV-infected cats, monocyte-associated FCoV viraemia was detected in all cats. At the end of the experiment, six cats still exhibited a monocyte-associated viraemia. Examination of the large intestinal content, collected at necropsy, demonstrated shedding of FCoV in seven cats, either by electron microscopy and/or by RT-PCR. Results are compiled in Table 3.

All animals had remained clinically healthy during the entire experiment.

Post-mortem analysis: FCoV viral load in different tissues

To determine to what extent FCoV was present in the organs of the cats, different tissues were analysed for presence of FCoV RNA.

In infection experiment I, the cats of group 2 were sacrificed at day 14 p.i., whereas those of groups 1 and 3 were sacrificed at day 28 p.i.

Table 3 Summary of the results obtained in the blood (antibody titres, monocyte-associated viraemia), in organs (FCoV-RT-PCR) and in large intestinal content (electron microscopy, FCoV-RT-PCR) of naturally FCoV-infected cats

Animal	TP1	TP2	TP3	TP4				Shedding			
	IFA	IFA	IFA	Monocytes	IFA	Monocytes	Spleen			Mes. Inn.	Bone marrow
1	≤25	800	200	+	50	+	+	-	+	-	+
2	≤25	≤25	200	+	≤25	(+)	n.p.	n.p.	n.p.	(+)	+
3	≤25	100	800	++	400	++	-	-	n.p.	+	n.p.
4	≤25	100	100	+	50	+++	+	-	-	-	+
5	≤25	400	400	++	400	-	-	+	+	-	+
6	≤25	>3200	≤25	++	50	++	+	-	-	-	-
7	≤25	>3200	400	+	400	-	+	+	-	+	+
8	≤25	≤25	100	+	≤25	(+)	-	-	-	+++	+

IFA: immune fluorescence analysis (titre); EM: electron microscopy (negative staining technique); mes.lnn.: mesenteric lymph nodes; n.p.: not performed. -: negative; (+): weak positive; +: moderate positive; ++: strong positive; +++: very strong positive. TP: time point. 1: directly prior to infection; 2: 2 months p.i.; 3: 5 months p.i.; 4: 7 months p.i. (euthanasia).

(14 days after re-infection), respectively. In all cats, macroscopical findings were restricted to enlargement of mesenteric lymph nodes. Samples of different organs (duodenum, jejunum, ileum, colon, mesenteric lymph nodes, liver, spleen, bone marrow, kidney, thymus, lung, tonsil and brain) were collected and examined for FCoV load. Ileum, colon and mesenteric lymph nodes displayed the highest viral loads (Fig. 1). In animals of group 2 (Fig. 1B), high viral loads were detected in duodenum, jejunum, liver, spleen and kidney. In all three groups, FCoV was detected in the entire intestine and in all organs tested, with the exception of the bone marrow.

In infection experiment II, the cats were sacrificed on day 17 p.i. Like in the first infection experiment, macroscopical findings were restricted to enlargement of mesenteric lymph nodes. The patterns of viral loads in the organs of these three animals were similar to those in the cats of group 2 of infection experiment I, infected with the same FCoV strain: duodenum, bone marrow and brain showed comparatively low viral loads, while jejunum, ileum, colon and mesenteric lymph nodes showed high viral loads (Fig. 2).

In infection experiment III, the animals were euthanased 80 days after infection. Macroscopical findings were again restricted to enlargement of mesenteric lymph nodes. In addition to the organs examined in the previous experiments, skin and skeletal muscle samples were analysed. In the animals, which received the lowest dose of virus (Fig. 3A), the virus was present at low levels in almost all organs. With increasing infection doses, higher viral loads were detected in the colon (Fig. 3B and C). However, due to the small number of cats ($n=2$ per group), the differences are not significant.

In naturally infected cats, like in experimentally infected animals, macroscopical findings were restricted to enlargement of mesenteric lymph nodes. Cats tested positive for FCoV in spleen, bone marrow and/or mesenteric lymph nodes. Five of the seven cats examined were FCoV-positive in at least one of the organs. In four animals, FCoV-RNA was detected in spleen and/or bone marrow, two cats were positive in the lymph nodes. Three animals were positive in two of the three organs tested. None tested positive in all organs analysed (Table 3).

Histopathological findings

In addition, organs were subjected to histological examination for evaluation of pathological findings.

In all cats, both experimentally and naturally infected, histopathological findings were restricted to a generalized B and T cell hyperplasia in spleen and mesenteric lymph nodes. The thymus exhibited no or minimal involution. The bone marrow showed moderate to high activity. All other organs were unaltered.

Discussion

In the present study we investigated both FCoV distribution and loads in faeces, blood and organs of cats experimentally infected with different FCoV type I field strains as well as of naturally FCoV-infected cats.

So far, in the majority of experimental infection trials, the investigators used a type II FCoV (Chalmers et al., 1993; Fiscus et al., 1987; Glansbeek et al., 2002; Pedersen et al., 1981a, 1984a; Pedersen and Floyd, 1985; Vennema et al.,

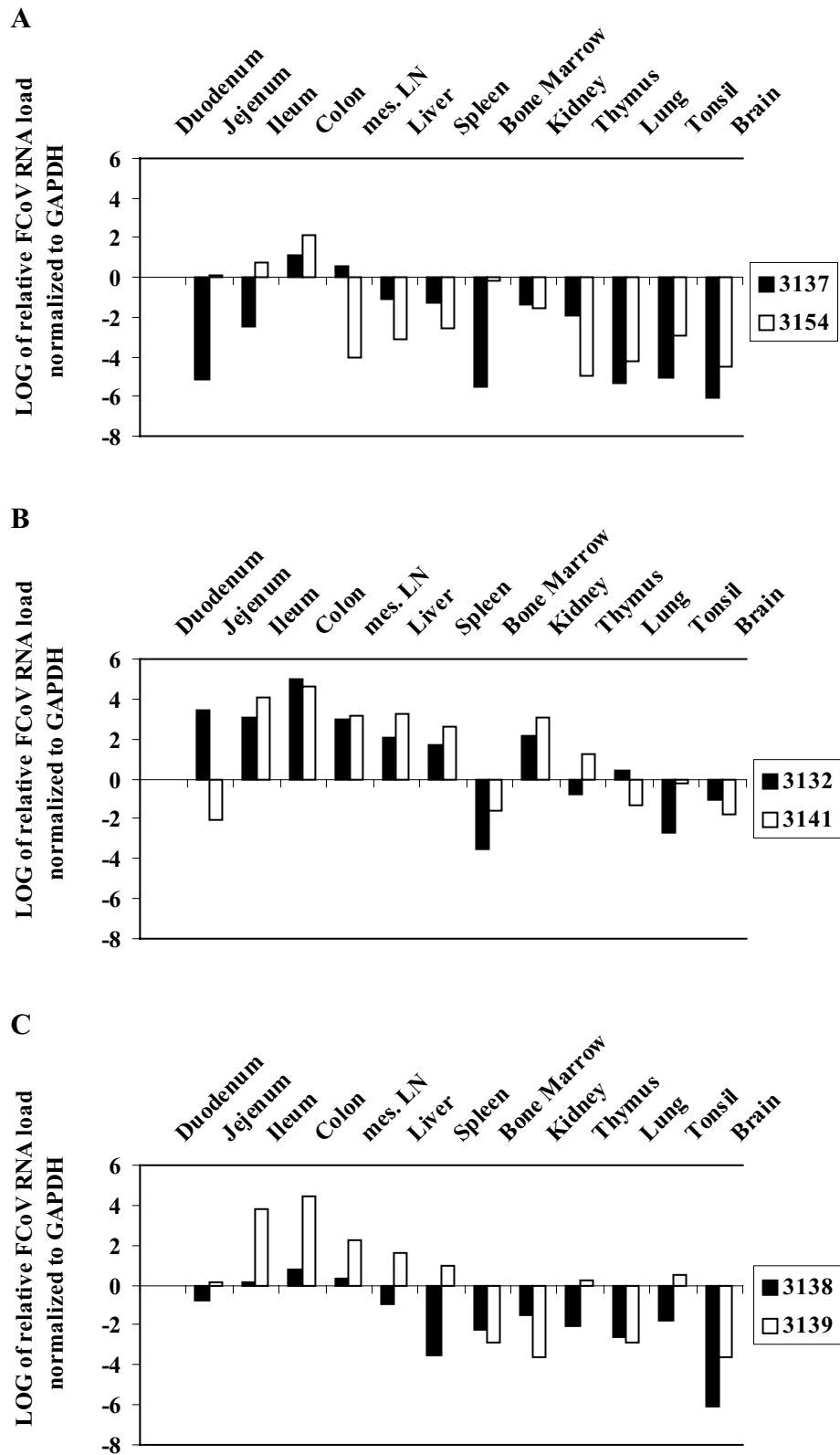


Figure 1 FCoV viral loads in organs of experimentally infected cats: Experiment I. Data are calculated using the comparative C_T method for relative quantitation. Expression in duodenum served as calibrator. A: group 1; B: group 2; C: group 3.

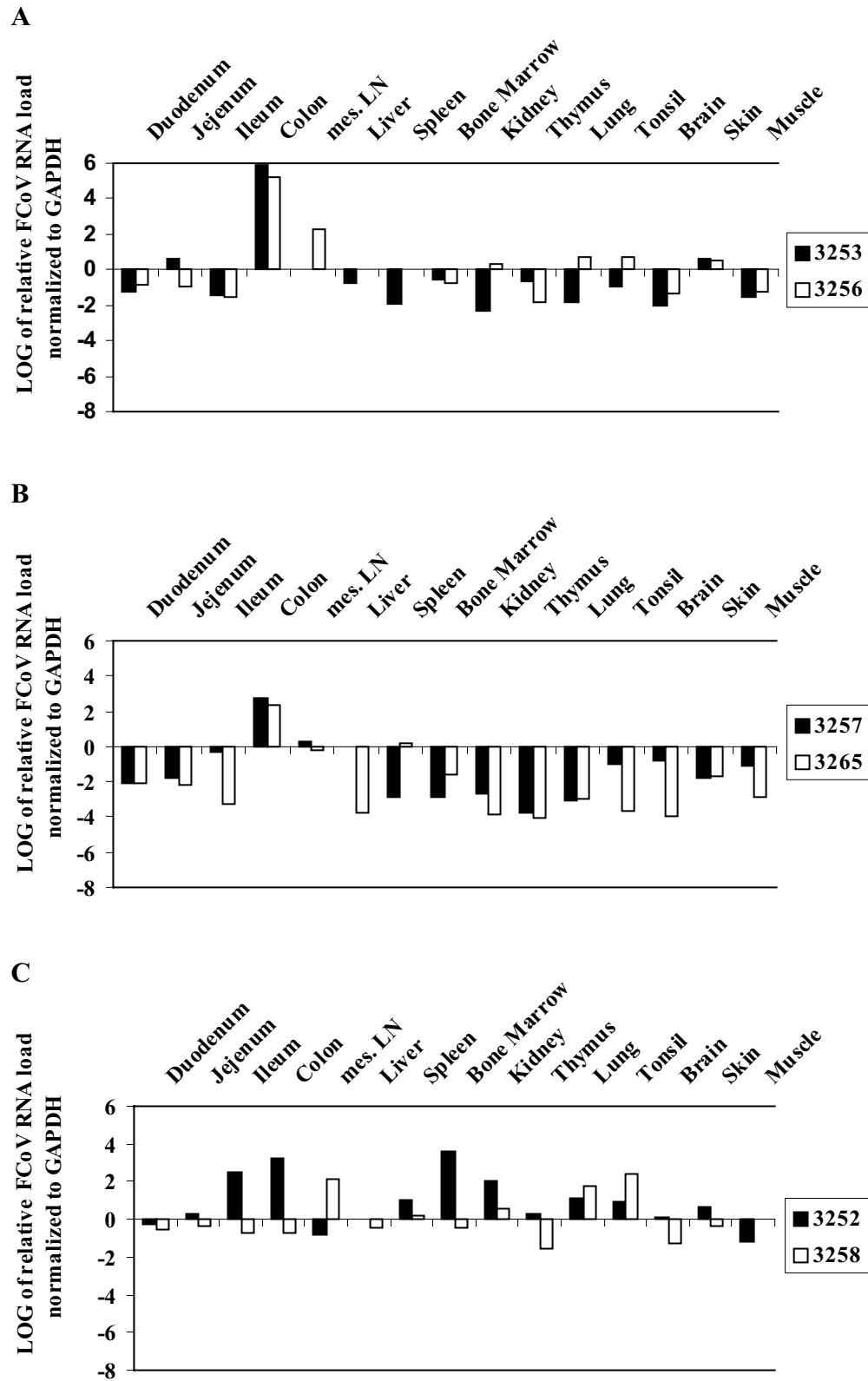


Figure 3 FCoV viral loads in organs of experimentally infected cats: Experiment III. Data are calculated using the comparative C_T method for relative quantitation. Expression in duodenum served as calibrator. A: group 5; B: group 6; C: group 7.

concluded that the high FCoV antibody titres observed in all eight animals represent a high FCoV load in all these cats (Gut, 1999).

In these naturally infected cats, 5 months after exposure to healthy FCoV-infected animals, blood monocytes were found to be positive for FCoV-RNA. Furthermore, at necropsy, monocytes were still positive in 75% of the cats. These findings are in accordance with other studies, which showed that FCoV-infected, healthy cats can maintain monocyte-associated viraemia for at least 12 months (Gunn-Moore et al., 1998). They differ from the results obtained from our experimentally infected cats where FCoV-RNA was only detected in the monocytes of one animal. However, this is likely due to the lower sensitivity of the FCoV RT-PCR, performed on RNA directly isolated from peripheral blood monocytes. In comparison, monocytes of naturally infected cats in this study as well as naturally infected cats in a previous study had been cultivated prior to FCoV RT-PCR (Gunn-Moore et al., 1998).

In addition, as in experimentally infected cats of this study, FCoV was found in the spleen, lymph nodes and/or bone marrow of most of the naturally infected cats at the time of necropsy. From these observations we can conclude that in cats infected with FCoV by natural contact with infected animals FCoV spreads systemically and into internal organs (probably via infected monocytes within the vasculature) (Kipar et al., 2003).

Several of the infected cats were found to be seronegative although they were shedding virus. This is not surprising and may be explained by the fact that in many cats, replication initially may be confined to the intestine and only later spread to the blood circulation and to the lymph nodes. As long as infection is confined to the intestine or the viral load is low, this may not induce seroconversion. Similar observations were also made in FIV infection in cats experimentally infected with low challenge doses (Boretti et al., 2000).

As FCoV could be detected in almost all organs tested from both experimental and naturally infected cats in the absence of any clinical signs and pathological findings, we concluded that development of FIP is not an imperative sequel of systemic spreading of FCoV. Originally, Pedersen and co-workers postulated that the apathogenic FCoV (designated FECV) remained restricted to the intestinal tract and that systemic infection would coincide with the development of FIP (Pedersen et al., 1981b). However, our observations support the hypothesis that the key event triggering the development of FIP is not associated with the capability

of an FCoV to cause viraemia and systemic infection. Although the observation period was not sufficiently long to see animals developing FIP, and we did not sequence the virus to detect mutations, these observations support the concept that FCoV mutants with increased virulence arise during replication (Kennedy et al., 2001; Vennema et al., 1998). To further substantiate this hypothesis it would be important to determine if FCoV is replicating in the blood. A recently developed method to determine FCoV messenger RNA in peripheral blood mononuclear cells could address this question (Simons et al., 2002).

In conclusion, we have demonstrated, that SPF cats experimentally infected by different FCoV type I field strains or naturally FCoV-infected, can remain clinically healthy despite overall high viral loads in faeces and different organs. It is suggested that identical mechanisms of FCoV kinetics occur under field conditions and that high viral loads are the basis for the generation of virulent FIPV strains.

Acknowledgements

The authors would like to acknowledge Armin Rüdumann for expert technical help with the experimentally infected cats as well as Edith Rhiner, Elizabeth Rogg and Beatrice Weibel for excellent technical support. Thanks are due to Dr W. Leukert for clinical supervision of the naturally infected cats. We are grateful to Mr A. Barth for perfect technical assistance.

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